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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

TECH CENTER 1600/2900

In re application of

Wolfgang M. Franz et al.

Serial No. 09/068,751

filed: November 14, 1996

for: Gene Therapeutic Nucleic Acid Working Model and Its Production and
Use for Treating Heart Disease

DECLARATION UNDER RULE 132

I, the undersigned, Wolfgang M. Franz, a citizen of Germany,
do hereby depose and declare:

I have graduated in medicine and I was awarded the degree of PhD from the
Technical University of Munich, Germany, in 1986.

My major research interest is directed to gene therapy of cardiovascular diseases.

I am co-inventor of the subject matter of U.S. Patent Application No. 09/068,751.

I have carefully read the Office Action dated December 20, 2001 and the
specification of the present patent application.

With regard to the Examiner's contention that the specification does not enable any
person skilled in the art to make and / or use the invention commensurate in scope
with the present claims I would like to provide the following response:

The present application is, to my opinion, the first document disclosing the successful
cardiac specific, in particular ventricle specific, gene transfer in neonatal or adult
animals by making use of viral vector systems.

In my opinion, a skilled person would have been enabled by the detailed disclosure of the specification to perform the invention, in particular the tissue specific gene transfer in adult or neonatal animals.

The experiments disclosed in the specification illustrate operability of a **functional** regulatory nucleic acid sequence as obtained from the **rat** *mlc-2* gene under conditions of somatic gene transfer (see in particular Examples 8 to 11 of the specification).

Meanwhile we were able to locate the corresponding regulatory sequence of the **human** *mlc-2* gene. The obtained nucleotide sequence is attached as

ANNEX I

We have analysed the sequence for potential regulatory elements and could locate sequence motifs **almost identical** to those disclosed in the present specification and considered as vital for the intended purpose. These elements are, in 5'-3' direction, the following: CSS-like sequence, MLE1-box, HF-1a-, HF1b-, HF-2-, HF-3- and E-box. The same sequence motifs have been identified in the specification to play a critical or at least favourable role in the regulation of cardiac specific gene expression.

Said high degree of similarity observed between rat and human regulatory elements is illustrated by the partial sequence alignment attached as

ANNEX II

The rat sequence motifs are shown below the corresponding human sequence motifs. Identical nucleotide residues are identified by a vertical line between both sequences.

On the basis of said additional experimental evidence, I am convinced that a skilled person will not have to practice "trial and error" experimentation in order to provide

viral constructs other than those exemplified in the specification which can be successfully used for cardiac specific, in particular ventricle specific, gene transfer.

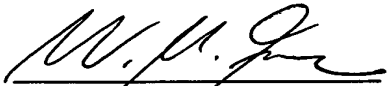
I would also expect that a similar favourable cardiac tissue specificity is also observed when the same regulatory elements are inserted in different viral vectors, because cardiac tissue specificity, which, in my opinion is the key for successful and valuable somatic gene transfer to the heart muscle, is mainly influenced by the cardiac tissue specificity of the promoter sequence operatively linked to the gene to be expressed with cardiac specificity.

In this respect, I would like to point to a comparative experiment already disclosed in the present specification. Example 11 describes the differences in cardiac tissue specificity and activity of gene expression observed on the one hand for a recombinant virus vector construct of the present invention, i.e. **Ad-mlcLuc**, and, on the other hand, for a virus construct carrying, **in the same virus, a different muscle derived promoter**, i.e. alpha-mhc. Said construct is designated **Ad-mhcLuc**. As illustrated by Figures 8A and 8B merely Ad-mlcLuc, i.e. the viral construct of the present invention, provides for a strong and cardiac specific gene expression while Ad-mhcLuc also directs non-specific gene expression. Significant levels of non-specific gene expression were observed for Ad-mhcLuc in kidney, spleen, liver, diaphragm, lung and intercostal muscle. Moreover, the construct of the invention was three to four times more active in the heart than Ad-mhcLuc. Said experimental evidence supports my point of view that a skilled reader of the present specification will in fact be able to practice, without the need of "trial and error" experimentation, the invention within the scope of the claimed recombinant vectors, carrying a regulatory mlc-2 gene fragment which must be functional, i.e. directs cardiac specific gene expression.

The undersigned declares further that all statements made herein on his own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that wilful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and

that such wilful false statements may jeopardise the validity of this application or any patent issuing thereon.

Munich, 15-06-2002

A handwritten signature in black ink, appearing to read 'W. M. Franz', written over a horizontal line.

(Wolfgang M. Franz)

CAGTCTTGAG TGATGCTGAA AGGAACCCCT GAAGTCTACA AAGAACCAAG	50
TCCTCCCTGG ATTCTCCAAT CCCAGGGCCT TGTCCCTGGT CTTGGGGGCT	100
CCCTGGGGCA ACACAACCCA TTGATGAGAG AGACTTTGGA TCTCTGGCTC	150
TCTCAACAGA CCCCAAGGCC TAGTCCACAC CCCACGTGCT CCACGTCCCA	200
GCAGCCACGT GGTTCATGC CCCATTCAGG CCGCCATTTC CCAGAATCCC	250
TAGACACAAT CTCACTTAAT CCTCCCAGCA GCCTTATGGA GGTGTGTGAT	300
CTCCCTTTTC CAGGTGAGGA AACAGGCCCG AGAGGGTGAG TGCCCTATTT	350
GACAACCCCT CTTGCTATCC AGCCAGAATG GTTCCTCTAG TACCCCTTCT	400
GGAGGCCTGG CTGTACAGGT GTCCCT <u>CAGG GACACACACC CCCACTCGAC</u>	450
<u>TCTGGGGGCC CAGCCCATCC TAATCCCCAC</u> CCCGGGGCTT CCCACCCCCC	500
ATCATACACT CTCACATCT TCTGTGGCTG CAACAACCTT TTCACTTGGC	550
CAGTTGGAGC TACTGACTGC TCACACAGGG TTTTAACGAA AATCTATGGT	600
GTGCCTATTA GCTAGGGAAA CATTTATTCT GGTGTTGTCA GAGAACCTTG	650
GACAGAAAAG CTCCTCTTGA TGTGTGCACT GCACATATGT GGATGCGTGT	700
ACATGCACGT GTCTGTGTGC CTCTATGCAT GTGCAGACGT GTTTTTGTCT	750
GTGCATGCAT GTGCCTACAC ACACACATGA ACACATCTTT TGTTATTAAA	800
GATCTGTCAG AAGAGTGTCC TGGGTAAGTC TAACCCATGT GGGACTGCAG	850

AGAAGAAAAA AACCCACACC TTTTTTTGTC ACAGCCATCA ATGGTCCTTG	900
GGTTTGTGTG CCCCCAAATT GAGATTATTT TTCCACCTGA GAAGGGGAGT	950
GAGTGATAGC TACCATTGTC CAGGTCTCAC CTCCTTTTAC CCTCTGGAAA	1000
ACCTAATAAG AAAAGTGATT TCTTTTTTTA AGCTCTGGAA AACTCCAGCC	1050
CCAGGGGGCC TTCTGTTTCT CAAAGCCTCC AAATTCTCCC TGCCTTGAGG	1100
TATGTGCTGT CCCCCTGCC TGGAGCCCCC TTTGCAGACT CTGCTTGAG	1150
GAGCCCACCT GCGCCCCTGT CTGAGGCTGT CACCTGGCCA CTGCCATGCC	1200
TCTGTCTCAT CCCTGCATGA GATCCGTCAC TGCCTGCAAC TGTCTGGGTT	1250
GTGCATTTGT TTACTTTCTC CTTGTCCATC TTCCCCTCGC ACTTACGCAC	1300
CTCAAGGGAA GGAATTTGT TGCTTTGCGC TGGGCTCGAT GAAGGGGAAT	1350
GAATGCTGGT TCAGCCATCA GCCCCGCACC CACACTACTG GGAGGGCAGA	1400
GGGACATTCT CTTCTTAGA GGTGTGGCCT CTGGCACTCA GGCTTGCCAC	1450
CCACGGACAC TAAATAACCA CAATGATTCC AAGCCCCGAG TTCTTGCTCC	1500
CTGAATCCCA AGGCTGTCTT TAAGGGCACA GGAAGATGGC CATCTTTTGT	1550
TGTTTTGGTT TAGTTTGGG TTTTTTGGT CTTTGTTTTT GAGATGGAGT	1600
CTTGCTCTGT CGCCCAGGCT GGAGTGCAAT GGCACGATCT CGGCTCACTG	1650
CAACCTCCGC CTCCTGGGTT CAAGCAATCC TCCTGCCTTA GCCTCCCAAG	1700

TAGCTGGGAG TACAGGCATG TGCCACAACG CCCAGCTAAT TTTTGTATTT	1750
TTAGTAGAGA TGGGGTTCTG TCATGTTGGC CAGACTGGTC TTGAACTCCT	1800
GACCTCAGGT GATCTACCCG CCTCGGCCTC CCAAAGTGCT AGGTGTGAGC	1850
CAACATGCCC GTCCTTTTTT TTTTTTTTTT TTTTTTTGAG TCAGAGTCTC	1900
ACTCTGTGCG CCAGGCTGGA GTGCAATGGC GCTATCTCGG CTCACTGCAA	1950
CCTCTGCCTC TCGGCCTCAA GCGATTCTCC TGCCTCAGTC TCCTGAGTAG	2000
CTGGGACTAC AGGCCCGCGC CACAACGCCT GGCTAATTTT TGTATTTTTA	2050
GTAGCGACAA GTTTCATCAT GTTGGCCAAG GTCGTCTTGA ACTCCTGACT	2100
CAAGTGATCC ACCCGCCTCA GCCTCTCAA GTGCTGGCAT TTCAGGTATA	2150
AGCCACTGCA CCCAGCAGGA AAGCTGTCTT CAGTAAAAGT ATTATATAAT	2200
GACACCTTGC ATTCTGAGAG CAGCTGCTGT TTTCAAGGCT CTTAAAGAGC	2250
CTGGACTCTG GAGACAAAGG GGCCTCCAGA GGGGTCCACG CCTAGCTCCA	2300
TCACTGTGTG ACCCTGGGCA GCTCACTTCG CCTCTCTGAG CTTTTGTTTC	2350
CGCATCTGTA AAATGGGGGC ATGGATGATG AGGTGGTCCC CACCCTCTAG	2400
GGTGGCTGGA AAATTATGTG TGGGAGCCAT GAGCACATAG TGTCCGGCAC	2450
GTGCCAGTGC TCAGTCAATG AGATTTGTCA TTTCTTCAGT CAACAAATAT	2500
TTATTTTGA GCTGCTGCTG TGTGCATCAT GAGCTGGGAG CTGGGGAGAC	2550

AGTCAGTGGT GAGGGAACT AAAGTGATCC CTGCCCTCTG AGCTGACGCT	2600
CCACAGATGC TGAAGAAAAT GAGTCAGTGC ACTGTGGGCA GTGTTCTGGGA	2650
CTGCCTCACG CTGTGCAGAG AAACAAAGAA GGGAGATCGG AGCGCAGGAG	2700
GTGCGTGGCT GTGTTATTTG TTTGTTTTGA GACAGGGTCT TGCTCTGTCA	2750
CCCAGGCTGG AGTGCAGTGG TGTGATCGTG GCTCACCACA GCCTCAACCT	2800
CCCGGGCCCA AGTGATCCTC CTGTCCCAGC CTCCTGAATA GCTGAGACTA	2850
TAGGCATGCA CCACCACGCA CAGCTATTTT TTTTCTTTT GCGTAGAGAC	2900
AGGCATCTCC CTATGTCACC CAGGCTGGTC GCAAACCTCCT AGGCTCAAGC	2950
AATCTTCCCG CCTCGGCCTC CCGCCGTGCT GGGATTTCAG GCATGAGCCA	3000
CAGTGCCAGC CTTATGGTT ATTTTAAAGA TGGTGGTCGG GGAGGCTTCA	3050
CTCAGGAGAT GACATATGAG CAAAGATGCA GTGAAGGAGG TGAAGGAAGG	3100
AGCCGTGCGA TGAAGGAGG AAAGACATTC CAGGTAGAGG GCACACAGGT	3150
GCAAAGACCC TGAGGCCAGA TCCAGGCTGA TAAACAGAG CATTTTAGCA	3200
GTCTCCTCTC CCTGCCATTT TTTTCTCAA AATTGACAAG CACAAGTGTC	3250
CCCGGCCCAA GCACCGCAGA GAGCGCGCAG CATCTCTCCC CGTGACCATG	3300
ACCCAGCTAC TGCCTCTTTA <u>ACCTTGAATG</u> CCTTTTGGG <u>GGCTCACGTG</u>	3350
<u>TCACCCAGTG</u> GCGAGTGAGC <u>CACCCTTACT</u> <u>TCAGAAGAAC</u> <u>GGCATGGGGT</u>	3400

HF-3 element

MLE1 element

HF-2-Box

GGGGGGGCCT TAGGTGGTGC CCGCCTCACC ^{E-Box} TATGACTGCC ^{HF-1a u.} AAAAGCGGTC 3450

HF-1b-Box
ATGGGGTTAT TTTAAACAT GGGGAGGAAG TATTTATTGT TCCTGGGCTG 3500

CAGAGAGCTG GCGGAGTGT GGAATTCTTC TCGGGAGGCA GTGCTGGGTC 3550

CTTTCCACCA TG

CSS-lik sequence

CAGGGACACACACCCCACTCGACTCTGGGGGCCAGCC-CATCCTAAT human
||||||| | ||| ||||| ||||| | ||| |||
CAGGGACAGAT-----CAC-----CTCTGTGGGGCCAGACTCATAGTAAC rat

HF-3- and MLE1 elements

CTCTT TAACCTTGAATGC CTTTTTGGGGGC TCACGTGTC-A CCCAG human
 |||||
 CTCTT TAACCTTGAAGGC ATTTTGGGT-C TCACGTGTCCA CCCAG rat
 HF-3-Box MLE1 Box

HF-2-Element, E-Box, HF-1a- and HF-1b- elements

TGAGCCACC CTTACTTCAGAAGAACGGC ATGGGGTGGGGGGGCCTTAGGTGGTG human
 |||||
 TGAACGGCT CTTACTTCAGAAGAACGGC ATGGGGTGGGGGGGC-TTAGGTGGCC rat
 HF-2-Box

CCCGCCTCACCTA TGACTG CCAAAA GCGGTCATG GGGTTATTTTTA human
 |||| | |||||
 TCTGCCTCACCTA CAACTG CCAAAA GTGGTCATG GGGTTATTTTTA rat
 E-Box HF-1a-Box HF-1b-Box